

Figure S1. Comparison of genomic AT content between ssDNA (black squares, black line) and dsDNA (blue diamonds, blue line) phage genomes and those of their hosts for phages previously analyzed by Xia and Yuen (2005) (left), and for phages added to GenBank since that analysis (right).

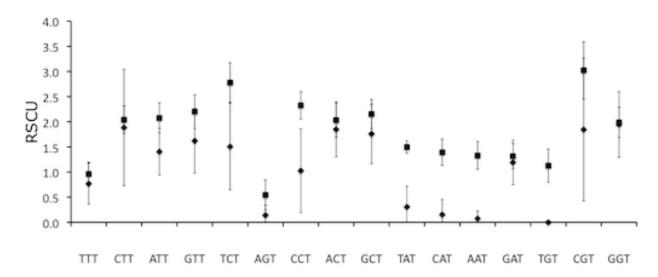


Figure S2. Mean RSCU values and 95% confidence intervals for NNT codons in *Microviridae* (squares) and *Inoviridae* (diamonds).